

SEQUENCE LISTING

<110> BERNSTEIN, Harold S.
COUGHLIN, Shaun R.

<120> METHODS AND COMPOSITIONS FOR REGULATING CELL CYCLE
PROGRESSION

<130> UCSF-020/02US

<140> Not Yet Available

<141> 2001-01-08

<150> US 09/156,316

<151> 1998-09-18

<150> US 60/060,688

<151> 1997-09-22

<160> 46

<170> PatentIn Ver. 2.1

<210> 1

<211> 802

<212> PRT

<213> Homo sapiens

<400> 1

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Glu	Ile	Leu	Lys	Ala	Ala	Val	Met	Lys	Tyr	Gly	Lys	Asn	Gln	Trp	Ser
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Arg	Ile	Ala	Ser	Leu	Leu	His	Arg	Lys	Ser	Ala	Lys	Gln	Cys	Lys	Ala
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Arg	Trp	Tyr	Glu	Trp	Leu	Asp	Pro	Ser	Ile	Lys	Lys	Thr	Glu	Trp	Ser
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Arg	Glu	Glu	Glu	Glu	Lys	Leu	Leu	His	Leu	Ala	Lys	Leu	Met	Pro	Thr
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Gln	Trp	Arg	Thr	Ile	Ala	Pro	Ile	Ile	Gly	Arg	Thr	Ala	Ala	Gln	Cys
			85						90					95	

Leu	Glu	His	Tyr	Glu	Phe	Leu	Leu	Asp	Lys	Ala	Ala	Gln	Arg	Asp	Asn
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Glu	Glu	Glu	Thr	Thr	Asp	Asp	Pro	Arg	Lys	Leu	Lys	Pro	Gly	Glu	Ile
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Asp	Pro	Asn	Pro	Glu	Thr	Lys	Pro	Ala	Arg	Pro	Asp	Pro	Ile	Asp	Met
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Asp	Glu	Asp	Glu	Leu	Glu	Met	Leu	Ser	Glu	Ala	Arg	Ala	Arg	Leu	Ala
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450				455				460							
Val 465	Lys	Gln	Met	Glu	Arg 470	Glu	Ser	Arg	Glu	His 475	Leu	Arg	Leu	Gly	Leu 480
Leu	Gly	Leu	Pro	Ala 485	Pro	Lys	Asn	Asp	Phe 490	Glu	Ile	Val	Leu	Pro	Glu 495
Asn	Ala	Glu	Lys 500	Glu	Leu	Glu	Glu	Arg 505	Glu	Ile	Asp	Asp	Thr 510	Tyr	Ile
Glu	Asp	Ala 515	Ala	Asp	Val	Asp	Ala 520	Arg	Lys	Gln	Ala	Ile 525	Arg	Asp	Ala
Glu 530	Arg	Val	Lys	Glu	Met	Lys 535	Arg	Met	His	Lys	Ala 540	Val	Gln	Lys	Asp
Leu 545	Pro	Arg	Pro	Ser	Glu 550	Val	Asn	Thr	Glu	Ile 555	Leu	Arg	Pro	Leu	Asn 560
Val	Glu	Pro	Pro	Leu 565	Thr	Asp	Leu	Gln	Lys 570	Ser	Glu	Glu	Leu	Ile 575	Lys
Lys	Glu	Met	Ile 580	Thr	Met	Leu	His	Tyr 585	Asp	Leu	Leu	His	His 590	Pro	Tyr
Glu	Pro	Ser 595	Gly	Asn	Lys	Lys	Gly 600	Lys	Thr	Val	Gly	Phe 605	Gly	Thr	Asn
Asn 610	Ser	Glu	His	Ile	Thr	Tyr 615	Leu	Glu	His	Asn	Pro 620	Tyr	Glu	Lys	Phe
Ser 625	Lys	Glu	Glu	Leu	Lys 630	Lys	Ala	Gln	Asp	Val 635	Leu	Val	Gln	Glu	Met 640
Glu	Val	Val	Lys	Gln 645	Gly	Met	Ser	His	Gly 650	Glu	Leu	Ser	Ser	Glu 655	Ala
Tyr	Asn	Gln 660	Val	Trp	Glu	Glu	Cys	Tyr 665	Ser	Gln	Val	Leu	Tyr 670	Leu	Pro
Gly	Gln	Ser 675	Arg	Tyr	Thr	Arg	Ala 680	Asn	Leu	Ala	Ser	Lys 685	Lys	Asp	Arg
Ile 690	Glu	Ser	Leu	Glu	Lys	Arg 695	Leu	Glu	Ile	Asn	Arg 700	Gly	His	Met	Thr
Thr 705	Glu	Ala	Lys	Arg	Ala 710	Ala	Lys	Met	Glu	Lys 715	Lys	Met	Lys	Ile	Leu 720
Leu	Gly	Gly	Tyr	Gln 725	Ser	Arg	Ala	Met	Gly 730	Leu	Met	Lys	Gln	Leu 735	Asn
Asp	Leu	Trp 740	Asp	Gln	Ile	Glu	Gln 745	Ala	His	Leu	Glu	Leu	Arg 750	Thr	Phe
Glu	Glu	Leu	Lys	Lys	His	Glu	Asp	Ser	Ala	Ile	Pro	Arg	Arg	Leu	Glu

1	5	10	15
Gly Val Thr Pro Gln Arg Gln Val Val Gln Thr Pro Asn Thr Val Leu	20	25	30
Ser Thr Pro Phe Arg Thr Pro Ser Asn Gly Ala Glu Gly Leu Thr Pro	35	40	45
Arg Ser Gly Thr Thr Pro Lys Pro Val Ile Asn Ser Thr Pro Gly Arg	50	55	60
Thr Pro Leu Arg Asp Lys Leu Asn Ile Asn Pro Glu Asp Gly Met Ala	65	70	75
Asp Tyr Ser Asp Pro Ser Tyr Val Lys Gln Met Glu Arg Glu Ser Arg	85	90	95
Glu His Leu Arg Leu Gly Leu Leu Gly Leu Pro Ala Pro Lys Asn Asp	100	105	110
Phe Glu Ile Val Leu Pro Glu Asn Ala Glu Lys	115	120	

<210> 8

<211> 107

<212> PRT

<213> Schizosaccharomyces pombe

<400> 8

Ser Val Thr Ile Glu Val Arg Asn Gln Leu Met Asn Arg Glu Gln Ser	1	5	10	15
Ser Leu Leu Gly Gln Glu Ser Ile Pro Leu Gln Pro Gly Gly Thr Gly	20	25	30	
Tyr Thr Gly Val Thr Pro Ser His Ala Ala Asn Gly Ser Ala Leu Ala	35	40	45	
Ala Pro Gln Ala Thr Pro Phe Arg Thr Pro Arg Asp Thr Phe Ser Ile	50	55	60	
Asn Ala Ala Ala Glu Arg Ala Gly Arg Leu Ala Ser Glu Arg Glu Asn	65	70	75	80
Lys Ile Arg Leu Lys Ala Leu Arg Glu Leu Leu Ala Lys Leu Pro Lys	85	90	95	
Pro Lys Asn Asp Tyr Glu Leu Met Glu Pro Arg	100	105		

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<211> 119

<212> PRT

<213> Homo sapiens

<210> 11
 <211> 2837
 <212> DNA
 <213> Homo sapiens

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 aggggggct atggaggaat accgaggatg aaattctgaa agcagcggta atgaaatatg 180
 ggaaaaatca gtggtctagg attgcctcat tgctgcatag aaaatcagca aagcagtgcg 240
 aagccagatg gtatgaatgg ctggatccaa gcattaagaa gacagaatgg tccagagaag 300
 aagaggaaaa actcttgac ttggccaagt tgatgccaac tcagtggagg accattgctc 360
 caatcattgg aagaacagcg gccagtgct tagaacacta tgaatttctt ctggataaag 420
 ctgcccaaag agacaatgaa gaggaacaa cagatgatcc acgaaaactt aaacctggag 480
 aaatagatcc aaatccagaa acaaaaccag cgcgccctga tccaattgat atggatgagg 540
 atgaacttga gatgctttct gaagccagag cccgcttggc taatactcag ggaaagaagg 600
 ccaagaggaa agcaagagag aaacaattgg aagaagcaag acgtcttgct gccctccaaa 660
 aaagaagaga acttcgagca gctggcatag aaattcagaa gaaaagaaaa aggaagagag 720
 gagttgatta taatgccgaa atcccatttg aaaaaagcc tgcccttggg ttttatgata 780
 cttctgagga aaactaccaa gctcttgacg cagatttcag gaaattaaga caacaggatc 840
 ttgatgggga gctaagatct gaaaaagaag gaagagatag aaaaaagac aaacagcatt 900
 tgaaaaggaa aaaagaatct gatttaccat cagctattct tcaaactagt ggtgtttctg 960
 aatttactaa aaagagaagc aaactagtac ttccctgccc tcagatttca gatgcagaac 1020
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 ctggcataac aaattctgct tccagtacac ttttgtctga gtacaatgtc accaacaaca 1140
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 gatatcgatc ttacacattc tgtgtataaa gaccttaact ccacaggacg gacattttag 2700
 agtttaaatt attaaggcta tcattctttt agtaatgtca tatttgcaaa cttttttagt 2760
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 aaaacaaaat ataaaaa 2837

<210> 12
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<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: epitope for M2
monoclonal antibody

<400> 12
Asp Tyr Lys Asp Asp Asp Lys
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<210> 13
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 13
gatttaacat aa 12

<210> 14
<211> 9
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 14
ttaacataa 9

<210> 15
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<212> DNA
<213> Homo sapiens

<400> 15
aataaaatca aaatt 15

<210> 16
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<400> 16
aaaggggaac acttt 15

<210> 17
<211> 55
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<220>
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<220>
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 <223> n = Any Nucleotide

<400> 17
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<210> 18
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<220>
 <223> Description of Artificial Sequence: Synthetic

<400> 18
 gcgtcgacaa gctttctaga 20

<210> 19
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<220>
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<400> 19
 cgctcgaggg atccgaattc 20

<210> 20
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 <213> Artificial Sequence

<220>
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<400> 20
 atttaacata a 11

<210> 21
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic

<400> 21
 tatttaacat aa 12

<210> 22
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 <212> DNA
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<220>
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<400> 22
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12

<210> 23
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 23
gagttaacat aa

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<210> 24
<211> 12
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 24
gatgtaacat aa

12

<210> 25
<211> 12
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 25
gattgaacat aa

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<210> 26
<211> 12
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 26
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12

<210> 27
<211> 12
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 27
gatttaccat aa 12

<210> 28
<211> 12
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 28
gatttaatat aa 12

<210> 29
<211> 12
<212> DNA
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<220>
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<400> 29
gatttaacct aa 12

<210> 30
<211> 12
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 30
gatttaacag aa 12

<210> 31
<211> 12
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 31
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<210> 32
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12

<210> 40

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gtgttattga aa

12

<210> 41

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acccacgtct at

12

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<223> Description of Artificial Sequence: Synthetic

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12

<210> 43

<211> 12

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic

<400> 43

gttgagtagt at

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<210> 44
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<220>
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<210> 45
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<210> 46
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<220>
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ttaacataag atttaacata aactctagag 90

<210> 47
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<220>
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<400> 47
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<210> 48
<211> 120
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 48
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<210> 49
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<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic

<400> 49
gatataacat at 12

<210> 50
<211> 12
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<220>
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<400> 50
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